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Original Paper

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Associating sporadic, foodborne illness caused by Shiga toxin-producing *Escherichia coli* with specific foods: a systematic review and meta-analysis of case-control studies

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Abstract

Shiga toxin-producing *Escherichia coli* (STEC) infections are a significant public health issue, with foodborne transmission causing >1 million illnesses worldwide each year. We conducted a systematic review and meta-analysis (PROSPERO registry # CRD42017074239), to determine the relative association of different food types with sporadic illnesses caused by STEC. Searches were conducted from 01 August to 30 September 2017, using bibliographic and grey literature databases, websites and expert consultation. We identified 22 case-control studies of sporadic STEC infection in humans, from 10 countries within four World Health Organization subregions, from 1985 to 2012. We extracted data from 21 studies, for 237 individual measures in 11 food categories and across three status types (raw or undercooked, not raw and unknown). Beef was the most significant food item associated with STEC illness in the Americas and Europe, but in the Western Pacific region, chicken was most significant. These findings were not significantly moderated by the raw or cooked status of the food item, nor the publication year of the study. Data from the African, South-East Asian and Eastern Mediterranean subregions were lacking and it is unclear whether our results are relevant to these regions.

Introduction

Shiga toxin-producing *Escherichia coli* (STEC) infections are a significant public health issue worldwide [1]. Circa 2010, STEC infections transmitted via food caused more than 1 million illnesses, 128 deaths and nearly 13 000 Disability Adjusted Life Years [2]. Cost effective intervention to prevent such infections requires identifying the foods that are the most important vehicles of exposure.

To determine the specific food types associated with foodborne illnesses, different methods are used to investigate outbreaks vs. sporadic illnesses. In outbreak investigations, the goal is to identify the specific food exposure common across the cases and both retrospective cohort and case-control studies are used to meet this objective. For sporadic cases of illness, risk factors, including food types, are most commonly identified via case-control studies. In case-control studies, the association of cases with various food exposures can be quantified, typically through odds ratios (ORs) and meta-analyses of these studies may yield summary estimates for food exposures of interest [3]. Therefore, to inform future preventative action, the aim of this study was to determine the food types with the greatest association with sporadic STEC illness.

Methods

Review question/scope

The specific question addressed by this systematic review and meta-analysis was: what is the relative association of different foods with sporadic STEC illness? A search of the PROSPERO Registry, the Cochrane Library and PubMed revealed one potentially relevant systematic review and meta-analysis that examined the relative contribution of routes of exposure to STEC infection [4]. Because this review: assessed broader routes of transmission (e.g. food, person to person); did not assess specific foods (other than raw/under-cooked meat); and only included larger ($n \geq 20$) studies of multiple designs (including but not limited to case-control studies),

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we determined that a new systematic review with a more in-depth analysis of different food categories was needed. The protocol for this review is registered in PROSPERO (# CRD42017074239; https://www.crd.york.ac.uk/prospero/display_record.php?RecordID=74239) and PRISMA guidelines were followed (except where formatting conflicted with journal requirements).

Eligibility criteria

The PECOS framework, as defined by Sargeant and O'Connor [5], was used to define the review eligibility criteria, as follows. The *population* was all human populations, with no limitations by age or other participant characteristics, location, or context/settings. The *exposures* were all foods (e.g. hamburger, leafy greens); we neither considered drinking water (tap, bottled, or other), breastfeeding, nor nasogastric feeding as foods, nor did we include studies that assessed general nutrition (including malnutrition) as a risk factor for STEC infection. The *comparator* group was individuals who are not ill with STEC infection (i.e. controls; determined either via laboratory testing or by the absence of symptoms) and the *outcome* was sporadic illness caused by laboratory-confirmed STEC infection. The *study design* was case-control studies and thus the effect measure of interest was the OR.

Search strategy

The search strategy was developed in consultation with a medical librarian and was reviewed by an expert in systematic reviews of foodborne disease, who was not involved in the original strategy development (IY). The search terms were developed through Medline Ovid and then adjusted as needed for each individual database searched. Details about search term development are given in the Supplementary Materials (Section A). A list of the final search terms is available via PROSPERO. The search was not limited by language, location, study period, or any other characteristics.

Searches were conducted from 01 August to 30 September 2017, in the following bibliographic databases: Medline (OVID), EMBASE, Scopus, CAB Direct, African Journals Online, Asia Journals Online and Latin America Journals Online. We also searched: the European Food Safety Authority (EFSA) journal; five databases of grey literature sources (ProQuest Dissertations and Theses, E-Theses Online Services (ETHOS), OpenGrey, Agricultural Research Service and Current Research Information System); the main World Health Organization (WHO) website, the six WHO regional websites, the Food and Agriculture Organization of the United Nations (FAO) website and the Africa Centers for Disease Control and Prevention website. To identify any unpublished or pre-publication studies, we consulted with: authors from Hooman *et al.* [6] and Paudyal *et al.* [7]; WHO advisors from STEC-related reports identified on WHO websites; WHO regional public health contacts and members of the Joint FAO/WHO Core Expert Group on STEC/VTEC. Finally, we searched citation lists of all types of review articles on STEC identified during the search and the citation lists of our final set of references.

Citation collection, deduplication and screening

Citations were collected, managed, de-duplicated and screened in RefWorks (ProQuest LLC, 2017). Attempts were made to obtain English translations of articles in other languages; if suitable

translation could not be obtained, the title and abstract were put through Google Translate for relevance screening and, if relevant, the entire article was reviewed and extracted by a native speaker of the article's original language with expertise in epidemiology and a familiarity with foodborne disease.

For initial relevance screening, titles and abstracts were screened by two independent reviewers per reference with a third reviewer to resolve conflicts. Inclusion criteria for relevance screening were the study is about STEC; a case-control study; done in humans; not an outbreak investigation. Citations fulfilling these criteria, or with insufficient information, were advanced to full-text screening, which was also completed by two independent reviewers per reference (with a third to resolve conflicts), using standardised instructions. Inclusion criteria for full-text screening, in addition to those above, were: the study investigates the exposures (or risk factors) experienced by a series of cases, compared to the exposures (or risk factors) experienced by a series of controls; the controls are not cases of some other disease; cases are individuals with illness caused by STEC; and, the study assessed food exposures. Studies that passed full-text screening advanced to data extraction. Further details about screening processes are given in the Supplementary Materials (Section A).

Data extraction and risk of bias assessment

Data were extracted by two reviewers, with a third reviewer to resolve conflicts, using standardised forms and instructions. A full list of the extracted variables, including how we assessed the appropriateness of the laboratory methods used, is given in the Supplementary Materials (Section A). Risk of bias was assessed using the Newcastle-Ottawa Quality Assessment Scale (http://www.ohri.ca/programs/clinical_epidemiology/nosgen.pdf), modified to address critical items for case-control studies using relevant questions from the RTI International – University of North Carolina at Chapel Hill Evidence-based Practice Center Item Bank and its modifications [8,9]. Overall study quality was captured using the RTI Overall Assessment question ('are the results of the study believable taking study limitations into consideration?'), as well as a modified version of ROBINS-I, relevant to case-control studies [10]. Finally, we assumed that age was the most important confounder given the established relationship between age and the risk of STEC infection [11e.g.]. Thus, we assessed whether age was adequately controlled as part of our quality assessment by assessing the combined impact of the study's design and analyses.

Analysis

Data were stratified into different categories for analysis. Study countries were classified into WHO subregions [2]. Food items were categorised using the United States' Interagency Food Safety Analytics Collaboration's hierarchical categorisation scheme of mutually exclusive food categories [12]. In our registered protocol, we stated that raw/undercooked foods would be treated as separate items than cooked foods. However, given that many of the food items were reported with unknown raw/cooked status, we chose instead to group raw and cooked food items (e.g. categorize raw beef, cooked beef and beef of unknown status all as 'beef') and explore the impact of raw/cooked/unknown status in the meta-regression.

Descriptive analyses were conducted to summarise study characteristics. To calculate the individual, study-specific ORs for each

food category for which results were reported, the following process was used. For all instances where the number of cases and controls who were either exposed or unexposed to a given food were reported in the paper directly, we used these exact values to calculate the OR and standard error (S.E.) post hoc. For the remaining instances where ORs were reported, we used the reported univariate OR and 95% CI to back-calculate the OR and S.E. To do this, we designed an optimisation process in which we fitted the log-transformed OR to a normal distribution and minimised the sum of squared differences between the observed and the fitted 95% CI. For comparison purposes, an alternate analytic approach was also applied, in which the reported univariate OR was used for all instances where such values were reported and the reported number of cases and controls who were either exposed or unexposed to a given food category was then used when ORs were not given. We used crude ORs and data, unadjusted for confounders because not all studies included adjusted ORs. All studies, regardless of characteristic, were included in the subsequent analyses.

Summary univariate ORs and their corresponding 95% CIs were calculated for each food category, both overall and by WHO subregion, using a random effects meta-analysis model, with restricted maximum likelihood used to weight studies. Heterogeneity was assessed using the I^2 statistic. Publication bias was assessed using the following: Begg and Mazumdar's rank correlation test [13] and Egger's regression test [14]. When significant publication bias was present, we used Duval and Tweedie's trim-and-fill method [15] to explore the impact on model estimates. For the food categories with significant overall associations, meta-regressions were conducted to explore heterogeneity by examining the relationship between single study characteristics (i.e. WHO subregion, publication year, study population age and raw/cooked status) and the ORs for food exposures. For both the summary ORs and the meta-regressions, the effect of clustering by study was explored in a sensitivity analysis to ensure that a study with several food exposures in the same food category did not have inflated influence on the estimates [16,17]. To this end, we fitted multilevel meta-analysis models using the study as a random effect. All analyses were carried out in R using the 'metafor' package [18].

Results

Numbers of citations identified

Results from the search, including the number of citations identified, are shown in Figure 1. The majority of the 411 full-text articles screened were in English, but 30 were in 13 other languages (Japanese, $n = 9$; Spanish, $n = 7$; Portuguese, $n = 3$; French, $n = 2$; Czech, Chinese, Dutch, German, Hungarian, Italian, Romanian, Slovenian and Thai all $n = 1$). From these 411 articles, we identified 22 case-control studies of sporadic STEC infection in humans, from 10 countries within four WHO subregions (Region of the Americas A (AMR A) and B (AMR B), European Region A (EUR A) and the Western Pacific Region A (WPR A)), conducted from 1985 to 2012 (Table 1); study locations and timeframes are also shown in Figure 2. All 22 studies were published in English and were from the peer-reviewed, indexed literature [11,19,39–].

Three potentially relevant studies were identified beyond the peer-reviewed literature databases. Two potentially relevant doctoral theses [40,41] were identified in the grey literature search,

but full-text documents were unavailable. Expert consultation identified a case-control study of sporadic human non-O157 STEC infections in the USA (Dr Patricia Griffin, US Centers for Disease Control and Prevention, Atlanta, GA; personal communication), but the results were unavailable at the time of our analysis.

Description of the identified studies

In terms of risk of bias, all studies used an adequate case definition, but for three of the 22 studies (14%) [22,30,34], it was difficult to determine whether the cases were representative based on the information provided. Of the 22 studies, 12 (55%) excluded secondary cases of STEC, in one they were included and for nine studies, this information was not provided (Table 1). Of the 22 studies, 20 (91%) contained enough detail to demonstrate that the laboratory methods were adequate to identify STEC, one described the identification of presumptive STEC [35] and one did not provide adequate information to assess the laboratory methods used [25] (Table 1).

Only one study [25], published as a short report, did not provide an adequate description of control selection nor definition. All other studies used controls that were without symptoms of current gastrointestinal infection, rather than those with negative laboratory tests for STEC. Thirteen of the studies (59%) used different methods to identify cases vs. controls (Table 1); in all these studies, cases were identified via existing health system mechanisms, including laboratory-based surveillance, whereas controls were predominantly identified via random or semi-random sampling from the population. Considering feasibility, validity, ethical and other issues, control selection was considered appropriate in 21 of the 22 studies (96%; i.e. controls represent the population from which the cases arose and, if the controls had acquired STEC infection, they would have been included as cases in the study), with the exception of one study [25].

Assessing how studies controlled for age in the study design, analysis, or both, two of the 22 studies (9%) [19,22] did not appear to adequately control for age. Both these studies matched on age during control selection, but they did not account for this in their analysis. Of the 20 studies that adequately controlled for age, two (10%) did not match on age during control selection, but adequately adjusted for age by including it in their regression models [38,39] and the remaining 18 (90%) matched on age during control selection, as well as conducted analysis that accounted for matching on age.

In all studies, exposures were ascertained via interview using comparable questions for cases vs. controls. All studies assessed case exposures during the incubation period prior to illness, whereas for controls, half assessed control exposure during the window period prior to the control interview, while the other half assessed control exposure during the same calendar period as the cases (Table 2). Of the 10 studies that assessed control exposure during the same calendar period as the cases, only the three most recently published ones [32,34–] reported the time elapsed between the exposure window and the case/control interview dates. In two of the three studies [32,34] the time elapsed for cases and controls was comparable (≤ 4 days difference), whereas in one study [33], controls were interviewed a median of 3 weeks later than cases were, suggesting the potential for differential recall bias. Non-response rates for cases and controls and descriptions of non-respondents were not given in most studies (17/22, 77%; Table 2). In 20 of the 22 studies (91%), the statistical

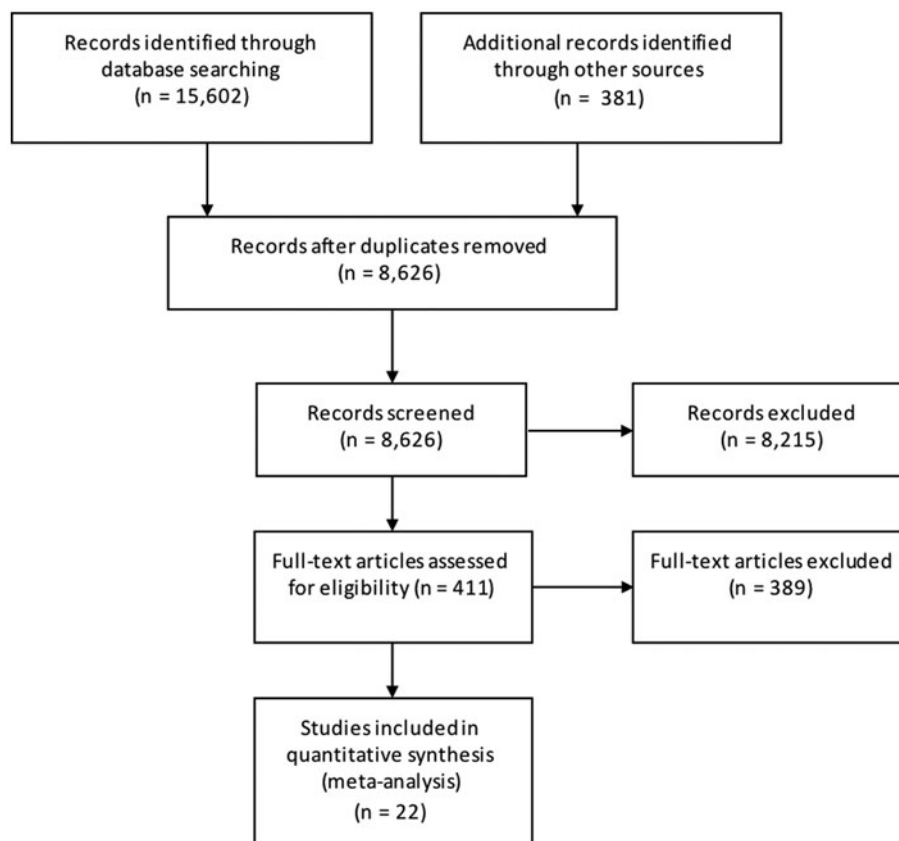


Fig. 1. PRISMA diagram showing the results of the search for case-control studies of sporadic STEC infections in humans (all dates and locations).

methods applied were considered adequate to determine ORs for food exposures; in one study, there was insufficient information to make this assessment [25] and in one study, the statistical methods were considered inadequate [22].

In considering all risk of bias assessment items together, two of the 22 studies (9%) were not considered reliable, taking study limitations into consideration [22,25]. Of the 22 studies, six (27%) were assessed to have a low risk of bias in the reported ORs for food exposures; 12 (55%) were assessed to have a moderate risk of bias in the reported ORs for food exposures, with the bias likely towards the null (i.e. towards an OR = 1); three (14%) were assessed to have a serious risk of bias (either toward or away from the null); and one did not have adequate information to make an assessment (Table 2).

Of the 22 studies, 18 (82%; Table 1) included individuals of all ages, with 15 providing results for all age groups combined in their estimates, two providing results stratified by age [11,38] and one providing results for both all participants combined and for the subset of children [30]. Of the 22 studies, four (18%) included only children. Most studies (16/22; 73%) included cases and controls drawn from the general population, with cases identified either via existing laboratory surveillance with public health notification (10/16; 63%) or via active case ascertainment at the laboratory level (6/16; 38%) and controls identified via random/semi-random sampling of the general population (including via existing registries, control databases, or random digit dialing; 8/16; 50%), or via the same facility or practice (5/16; 31%) or the same neighbourhood as the case (2/16; 13%). The remaining six of the 22 studies (27%) drew cases from specific facilities, via active case ascertainment within laboratories (3/6; 50%), emergency rooms (1/6; 17%) and by physicians (1/6; 17%), or health

record reviews (1/6; 17%). These six studies selected controls from the same facility as the cases (4/6), as well as from the case's friends (1/6) and neighbourhood (1/6). In 20 of the 22 studies (91%), cases were defined as symptomatic individuals with laboratory confirmation of STEC. In one study, cases were defined as those with post-diarrhoeal hemolytic-uremic syndrome, of whom 82.4% had a laboratory-confirmed STEC infection [22]. In another study, cases were either symptomatic individuals with laboratory confirmation of STEC or those with post-diarrhoeal hemolytic-uremic syndrome [34].

Food items associated with STEC infection

Extractable information on the relative odds of exposure to a given food, for cases as compared to controls, was provided by all but one paper [36]. Thus, we extracted data from 21 papers, for 245 individual measures in 11 food categories and across three status types: raw or undercooked, not raw (i.e. adequately cooked, treated, pasteurized, or other mechanism) and unknown (Table 3). Of the 245 individual measures extracted, 237 provided useable data (Table 4). Within the dairy category, the food items could not be divided by animal source because this information was only available for two items from one study (ewes' milk cheese and goats' milk cheese [33]). Similarly, the animal source was not provided for 'eggs', which were reported in two studies [31,37]. The 62 items classified as 'meat – unspecified' included items (of which 60 had useable data) that could not be assigned to their animal origin (e.g., beef, pork; Table 3). Of the 38 items classified as 'produce' only 11 were reported as specific fruits or vegetables (Table 3). Because there were very few results per specific produce item, this category was not divided further.

Table 1. Characteristics of the 22 case-control studies of non-outbreak (i.e. sporadic) STEC infection in humans, ordered by study timeframe (oldest to newest)

Lead author (Year published)	Country (WHO subregion ^a)	Study timeframe	Study Pop. Age	Study Pop. Type	No. cases; no. controls	Types of cases (all lab. confirmed?)	Secondary cases included/excluded	Case finding method	Control type	STEC category	Lab. methods adequate to identify STEC?
MacDonald (1988)	USA (AMR A)	1985–1986	All	Patients of specific facilities	24; 48	Non-outbreak GI illness with positive lab. result (yes)	Included (1 of the 24 cases was secondary)	Active case finding (at laboratory level)	Facility/ practice	O157	Yes (confirmed STEC O157:H7)
Bryant (1989)	Canada (AMR A)	1986–1987	All	Patients of specific facilities	81; 96	Non-outbreak GI illness with positive lab. result (yes)	Not described	Active case finding (in ER)	Friends	O157	Yes (confirmed STEC O157)
Le Saux (1993)	Canada (AMR A)	1990	All	General pop.	110; 220	Non-outbreak positive lab. result (yes)	Excluded	Active case finding (at laboratory level)	Neighbours	O157	Yes (confirmed STEC O157:H7)
Rowe (1993)	Canada (AMR A)	1990	Children (0–14 years)	Patients of specific facilities	34; 102	Post-diarrhoea cases of HUS (no; 88% were + for VTEC)	Not described	Active case finding (by physicians)	Facility/ practice	STEC	Yes (confirmed STEC)
Slutsker (1998)	USA (AMR A)	1990–1992	All	Patients of specific facilities	73; 142	Non-outbreak GI illness with positive lab. result (yes)	Not described	Active case finding (at laboratory level)	Facility/ practice	O157	Yes (confirmed STEC O157)
Holton (1999)	Canada (AMR A)	1991	All	General pop.	100; 200	Non-outbreak GI illness with positive lab. result (yes)	Not described	Lab.-based surveillance with public health notification	Neighbours	O157	Yes (confirmed STEC O157:H7)
Finelli (1995)	USA (AMR A)	1994	All	General pop.	23; 46	Non-outbreak GI illness with positive lab. result (yes)	Not described	Lab.-based surveillance with public health notification	Inadequately described	O157	Inadequately described
Mead (1997)	USA (AMR A)	1994	All	General pop.	22; 45	Non-outbreak GI illness with positive lab. result (yes)	Excluded	Active case finding (at laboratory level)	Population	O157	Yes (confirmed STEC O157:H7)
Parry (1998)	UK (EUR A)	1994–1996	All	General pop.	85; 142	Non-outbreak GI illness with positive lab. result (yes)	Excluded	Active case finding (at laboratory level)	Facility/ practice	O157	Yes (confirmed STEC O157)
O'Brien (2001)	UK (EUR A)	1996–1997	All	General pop.	369; 511	Non-outbreak GI illness with positive lab. result (yes)	Excluded	Active case finding (at laboratory level)	Facility/ practice	O157	Yes (confirmed STEC O157)
Kassenborg (2004)	USA (AMR A)	1996–1997	All	General pop.	196; 372	Non-outbreak GI illness with positive lab. result (yes)	Excluded	Active case finding (at laboratory level)	Population	O157	Yes (confirmed STEC O157)
Piérard (1999)	Belgium (EUR A)	Inadequately described; 1990's	All	Patients of specific facilities	37; 69	Non-outbreak GI illness, or HUS, with positive lab. result (yes)	Not described	Active case finding (at laboratory level)	Facility/practice	STEC	Yes (confirmed STEC)
Locking (2001)	UK (EUR A)	1996–1999	All	General pop.	183; 545	Non-outbreak GI illness, or HUS, with positive lab. result (yes)	Excluded	Lab.-based surveillance with public health notification	Facility/ practice	O157	Yes (confirmed STEC O157)

(Continued)

Table 1. (Continued.)

Lead author (Year published)	Country (WHO subregion ^a)	Study timeframe	Study Pop. Age	Study Pop. Type	No. cases; no. controls	Types of cases (all lab. confirmed?)	Secondary cases included/ excluded	Case finding method	Control type	STEC category	Lab. methods adequate to identify STEC?
Voetsch (2007)	USA (AMR A)	1999–2000	All	General pop.	283; 534	Non-outbreak GI illness with positive lab. result (yes)	Excluded	Active case finding (at laboratory level)	Population	O157	Yes (confirmed STEC O157)
Vaillant (2009)	France (EUR A)	2000–2001	Children (0–15 years)	General pop.	61; 114	Post-diarrhoeal HUS with confirmation of STEC (yes)	Not described	HUS surveillance with public health notification	Facility/ practice	STEC	Yes (confirmed STEC)
Rivas (2008)	Argentina (AMR B)	2001–2002	Children (0–15 years)	Patients of specific facilities	150; 299	Non-outbreak GI illness, or HUS, with positive lab. result (yes); also post-diarrhoeal HUS (no)	Not described	Health record review	Neighbours	O157	Yes (confirmed STEC O157)
Werber (2007)	Germany (EUR A)	2001–2003	All	General pop.	202; 202	Non-outbreak GI illness, or HUS, with positive lab. result (yes)	Not described	Lab.-based surveillance with public health notification	Population	STEC	Yes (confirmed STEC)
Hundy (2004)	Australia (WPR A)	2002	All	General pop.	11; 22	Non-outbreak illness/blood in stool with positive lab result (no)	Excluded	Lab.-based surveillance with public health notification	Population	STEC	Yes (presumptive STEC)
Denno (2009)	USA (AMR A)	2003–2005	Children (0–19 years)	General pop.	39; 75	Non-outbreak GI illness with positive lab. result (yes)	Excluded	Lab.-based surveillance with public health notification	Practice/ facility	O157	Yes (confirmed STEC O157)
McPherson (2009)	Australia (WPR A)	2003–2007	All	General pop.	113; 304	Non-outbreak GI illness with positive lab. result (yes)	Excluded	Lab.-based surveillance with public health notification	Population	STEC	Yes (confirmed STEC)
Friesema (2015)	The Netherlands (EUR A)	2008–2012	All	General pop.	208; 1563	Non-outbreak GI illness with positive lab. result (yes)	Excluded	Lab.-based surveillance with public health notification	Population	STEC	Yes (confirmed STEC)
Jaros (2013)	New Zealand (WPR A)	2011–2012	All	General pop.	113; 506	Non-outbreak GI illness, or HUS, with positive lab. result (yes)	Excluded	Lab.-based surveillance with public health notification	Population	STEC O157	Yes (confirmed STEC)

^aWHO Subregions comprise the following countries: Region of the Americas A (AMR A): Canada, Cuba, USA.

Region of the Americas B (AMR B): Antigua and Barbuda, Argentina, Bahamas, Barbados, Belize, Brazil, Chile, Colombia, Costa Rica, Dominica, Dominican Republic, El Salvador, Grenada, Guyana, Honduras, Jamaica, Mexico, Panama, Paraguay, Saint Kitts and Nevis, Saint Lucia, Saint Vincent and the Grenadines, Suriname, Trinidad and Tobago, Uruguay, Venezuela (Bolivarian Republic of).

European Region A (EUR A): Andorra, Austria, Belgium, Croatia, Cyprus, Czech Republic, Denmark, Finland, France, Germany, Greece, Iceland, Ireland, Israel, Italy, Luxembourg, Malta, Monaco, Netherlands, Norway, Portugal, San Marino, Slovenia, Spain, Sweden, Switzerland, United Kingdom of Great Britain and Northern Ireland.

Western Pacific Region A (WPR A): Australia, Brunei Darussalam, Japan, New Zealand, Singapore.

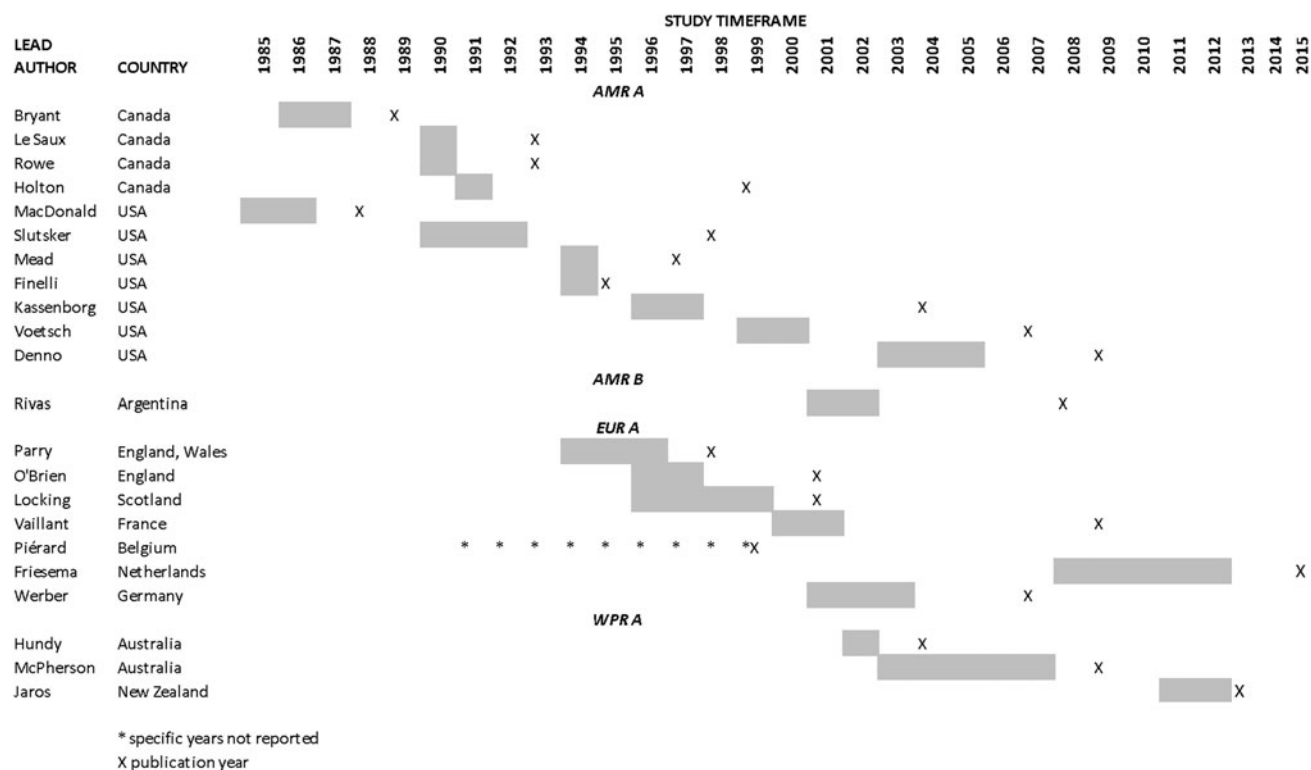


Fig. 2. Study locations and timeframes for the 22 identified case-control studies of sporadic STEC infections in humans.

Overall, beef and meat – unspecified were most significantly associated with sporadic STEC infection, although meat-unspecified became non-significant when the trim and fill method was used (Table 4; Fig. 3a and 3b). Produce, dairy, eggs and poultry/game-unspecified were also significant but had ORs less than one. When the alternate approach (see methods) to determining the ORs for each study/food was applied, estimates of the summary ORs neither changed in magnitude, nor direction, nor significance (Supplementary Materials, Section B). Under the sensitivity analysis, which accounted for clustering by study, only beef, eggs and poultry/game-unspecified remained significant.

Significant food categories varied moderately by WHO subregion (Table 5). In the Americas A region, beef and meat – unspecified remained the significant risk factors for STEC, whereas in the Americas B and European A regions, the only significant risk factor was beef and in Western Pacific A region the only significant risk factor was chicken. Under the alternate approach to determining the ORs for each study/food, our estimates of the summary ORs by WHO subregion changed: in the Americas A region beef became non-significant and in European A region meat-unspecified became significant (Supplementary Materials, Section B). Under the sensitivity analysis, beef remained the significant risk factor for STEC in the Americas A and B region and chicken in the Western Pacific A region; no factors remained significant risks in the European A region.

Our exploratory meta-regression analysis of the association between different study characteristics and ORs for food categories is shown in Table 6. Study population age was significant for dairy, as studies in children yielded lower ORs than studies of all ages. Study subregion was significant for meat – unspecified (with the study from the Americas B region yielding lower ORs than

studies from the Americas A region), produce (with studies from the European A region yielding higher ORs than studies from the Americas A region), dairy (with studies from the European A region yielding lower ORs than studies from the Americas A region) and chicken (with studies from the European A and Western Pacific A regions yielding higher ORs than studies from the Americas A region; data not shown). One of the two measures of risk of bias was significant for meat – unspecified, dairy and chicken, as follows. For both meat – unspecified and dairy, the study whose findings were considered not reliable yielded higher ORs than studies whose findings were considered believable (whereas findings did not differ significantly by Robin's I). For chicken, studies with serious risk of bias, either toward or away from the null, yielded lower ORs than studies with a low risk of bias (whereas findings did not differ significantly by overall study believability; data not shown). Publication year and whether the food item was raw/undercooked, not raw, or unknown were not significant moderating factors. Results under the alternate approach varied slightly and are given in the Supplementary Materials (Section C). Under the sensitivity analysis, even fewer significant modifiers remained.

Discussion

In this study, our aim was to determine the relative importance of different foods for sporadic, foodborne illnesses caused by STEC. Here, we found that beef was the most significant food item risk factor for STEC illness, significant in the Americas and Europe but not in the Western Pacific region, where chicken was the most significant food item risk factor. These findings for beef and chicken were not significantly moderated by the raw or cooked status of the food item, nor the publication year of

Table 2. Selected risk of bias assessment indicators for the 22 case-control studies of non-outbreak (i.e. sporadic) STEC infection in humans, ordered by study timeframe (oldest to newest)

Lead author (Year published)	Country (WHO subregion)	Study timeframe	Exposure window: cases	Exposure window: controls	Non-response rate and non-respondents	Robins- <i>I</i> ^a
MacDonald (1988)	USA (AMR A)	1985–1986	7 days before illness	7 days before interview	Not described	Serious
Bryant (1989)	Canada (AMR A)	1986–1987	7 days before illness	Same calendar dates as case	Same non-response rate for cases and controls	Serious
Le Saux (1993)	Canada (AMR A)	1990	10 days before illness	Same calendar dates as case	Not described	Moderate, towards null
Rowe (1993)	Canada (AMR A)	1990	14 days before illness	14 days before interview	Not described	Serious
Slutsker (1998)	USA (AMR A)	1990–1992	7 days before illness	7 days before interview	Not described	Moderate, towards null
Holton (1999)	Canada (AMR A)	1991	7 days before illness	Same calendar dates as case	Not described	Moderate, towards null
Finelli (1995)	USA (AMR A)	1994	7 days before illness	Not reported	Not described	No information
Mead (1997)	USA (AMR A)	1994	7 days before illness	Same calendar dates as case	Different non-response rates for cases versus controls, with non-respondents described	Moderate, towards null
Parry (1998)	UK (EUR A)	1994–1996	7 days before illness	Same calendar dates as case	Not described	Low
O'Brien (2001)	UK (EUR A)	1996–1997	5 days before illness	Same calendar dates as case	Different non-response rates for cases versus controls, with non-respondents not described	Low
Kassenborg (2004)	USA (AMR A)	1996–1997	5 days before illness	5 days before interview	Not described	Low
Piérard (1999)	Belgium (EUR A)	Inadequately described; 1990's	14 days before illness	14 days before interview	Not described	Moderate, towards null
Locking (2001)	UK (EUR A)	1996–1999	14 days before illness	Same calendar dates as case	Not described	Moderate, towards null
Voetsch (2007)	USA (AMR A)	1999–2000	7 days before illness	Same calendar dates as case	Not described	Low
Vaillant (2009)	France (EUR A)	2000–2001	7 days before illness	Same calendar dates as case	Not described	Moderate, towards null

(Continued)

Table 2. (Continued.)

Lead author (Year published)	Country (WHO subregion)	Study timeframe	Exposure window: cases	Exposure window: controls	Non-response rate and non-respondents	Robins- <i>I</i> ^a
Rivas (2008)	Argentina (AMR B)	2001–2002	7 days before illness	Same calendar dates as case	Not described	Moderate, towards null
Werber (2007)	Germany (EUR A)	2001–2003	10 days before illness	10 days before interview	Different non-response rates for cases versus controls, with non-respondents not described	Low
Hundy (2004)	Australia (WPR A)	2002	10 days before illness	10 days before interview	Not described	Moderate, towards null
Denno (2009)	USA (AMR A)	2003–2005	2–8 days before illness	2–8 days before interview	Not described	Low
McPherson (2009)	Australia (WPR A)	2003–2007	10 days before illness	10 days before interview	Not described	Moderate, towards null
Friesema (2015)	The Netherlands (EUR A)	2008–2012	7 days before illness	7 days before interview	Not described	Moderate, towards null
Jaros (2013)	New Zealand (WPR A)	2011–2012	14 days before illness	14 days before interview	Different non-response rates for cases versus controls, with non-respondents described	Moderate, towards null

^aModified ROBINS-I categories:

1. low risk of bias in the reported OR's for food exposures.
2. moderate risk of bias in the reported OR's for food exposures, with the bias likely towards the null (i.e. towards an OR = 1).
3. moderate risk of bias in the reported OR's for food exposures, with the bias likely away from the null (i.e. away from an OR = 1).
4. serious risk of bias (either toward or away from the null): the study has some important problems.
5. critical risk of bias (either toward or away from the null): the study is too problematic to provide useful evidence.
6. no information.

Table 3. Categories of the 245 food items extracted from the 21 case-control studies of non-outbreak (i.e. sporadic) STEC infection in humans, ranked in descending order by the number of food items per category

Food category (no. items within category; no. studies)	Types of foods items within the category (no. items; no. studies)	Number of items by cooked or processed status of the food item		
		Raw or undercooked	Not raw (i.e. cooked, treated, pasteurized)	Unknown/ not reported
Beef (83; 18)	Hamburger/ground beef (34; 15), beef (31; 5), beef sausage (4; 1), steak (3; 3), beef juice (3; 1), beef salami (3; 1), beef soup (2; 1), corned beef (2; 1), roast beef (1; 1)	35	1	47
Meat – unspecified (62; 13)	Meat (16; 7), deli meat (11; 5), minced meat (10; 2), sausage (8; 3), meat pies/empanadas (6; 1), hot dogs (3; 3), meatballs (3; 2), doner kebab (3; 1), meat casserole (1; 1), salami (1; 1)	14	10	38
Produce (38; 11)	Vegetables (14; 7), fruits/vegetables (6; 3), juice (5; 3), carrots (2; 1), fruit (1; 1), berries (1; 1), apricots (1; 1), cataloupe (1; 1), lettuce (1; 1), peaches (1; 1), plums (1; 1), strawberries (1; 1), tomatoes (1; 1), watercress (1; 1), watermelon (1; 1)	16	-	22
Dairy (25; 9)	Cheese (14; 4), milk (7; 4), milk/dairy products (2; 2), butter (1; 1), cream (1; 1)	11	2	12
Chicken (10; 8)	Chicken (9; 8), sliced processed chicken (1; 1)	1	1	8
Seafood (8; 4)	Fish (3; 2), shellfish (3; 2), seafood (1; 1), fish/seafood (1; 1)	-	-	8
Pork (7; 5)	Pork (4; 3), ham (3; 2)	-	-	7
Eggs (5; 2)	Eggs (5; 2)	-	-	5
Lamb (3; 1)	Lamb (3; 1)	-	-	3
Turkey (2; 2)	Turkey (2; 2)	-	-	2
Poultry/Game – unspecified (2; 2)	Poultry/game (1; 1), other poultry than chicken or turkey (1; 1)	-	-	2

Table 4. Results of the meta-analysis, showing pooled univariate odds ratios (ORs) per food category (significant values shown in bold), ranked in descending order by the number of food items in the category

Food Category (no. items within category; no. studies with useable data)	Odds ratio (95% confidence interval)	P-value	P-value Regression test	P-value Rank test	I ² (%)	Trim and Fill Method	
						Odds ratio (95% confidence interval)	P-value
Beef (80 ^a ; 18)	1.667 (1.408–1.975)^b	<0.001	<0.001	0.008	72	1.437 (1.205–1.713)	<0.001
Meat – unspecified (60 ^a ; 13)	1.281 (1.090–1.506)	0.003	<0.001	0.007	61	1.069 (0.894–1.279)	0.463
Produce (38; 11)	0.671 (0.534–0.843)	<0.001	0.035	0.119	72	0.671 (0.534–0.843)	<0.001
Dairy (23 ^a ; 9)	0.734 (0.558–0.966)	0.027	0.048	0.319	70	0.673 (0.500–0.906)	0.009
Chicken (9 ^a ; 8)	0.827 (0.377–1.814)	0.636	0.517	0.358	83	-	-
Seafood (8; 4)	0.758 (0.457–1.256)	0.282	0.902	0.905	75	-	-
Pork (7; 5)	1.032 (0.632–1.685)	0.900	0.201	0.239	63	-	-
Eggs (5; 2)	0.658 (0.515–0.841)^b	<0.001	0.504	0.483	0	-	-
Lamb (3; 1)	1.936 (0.582–6.441)	0.282	0.072	0.333	45	-	-
Turkey (2; 2)	1.055 (0.085–13.102)	0.967	N/A	1.000	65	-	-
Poultry/Game – unspecified (2; 2)	0.411 (0.228–0.740)^b	0.003	N/A	1.000	32	-	-

^aThese numbers are less than in Table 3 because some food items as reported did not have sufficient useable data^bThese items remained significant when clustering by study was accounted for

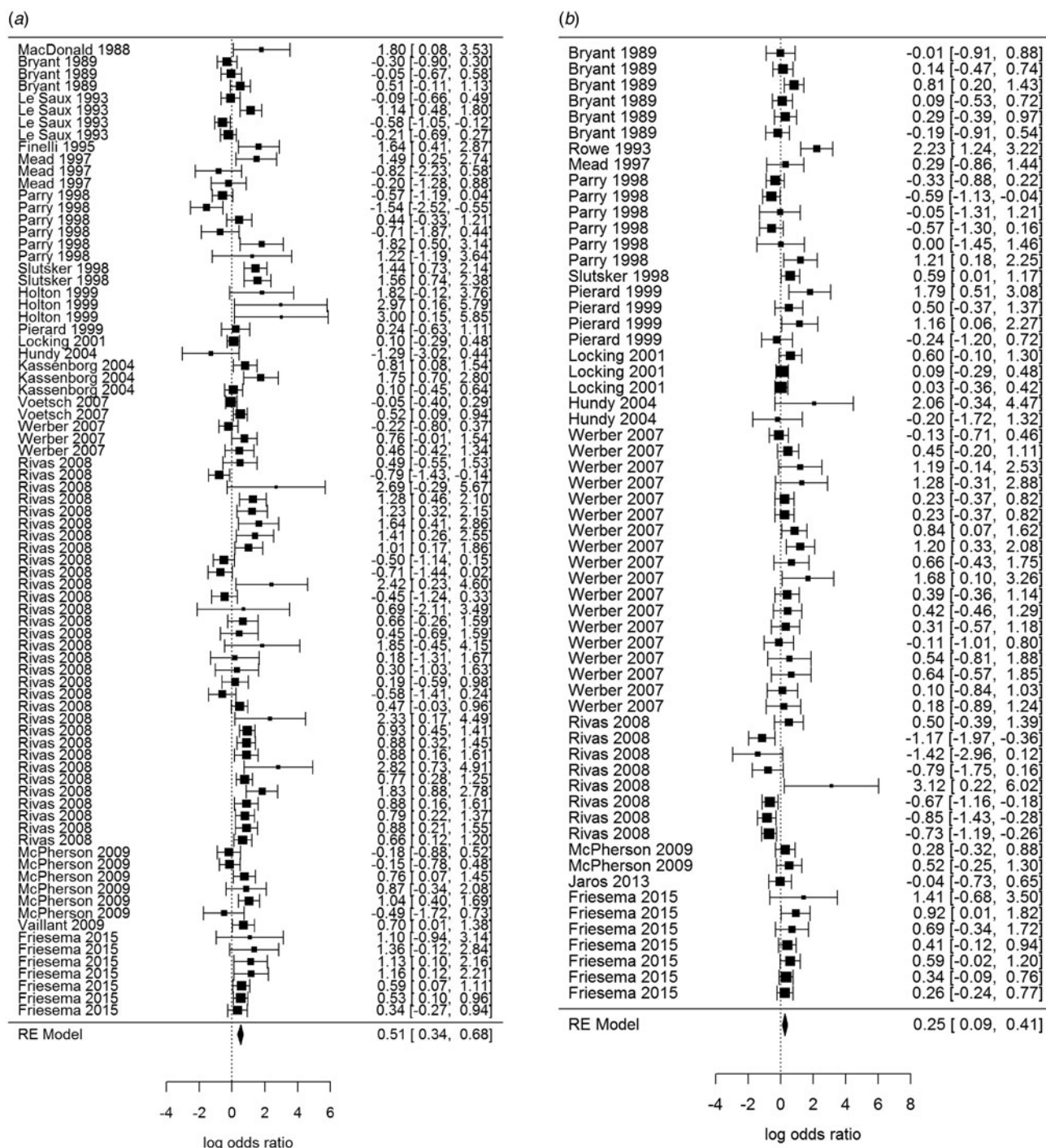


Fig. 3. Forest plots of the log odds ratio (OR) of the risk of human STEC infection from beef (a) and meat-unspecified (b), showing the overall pooled OR together with the 95% confidence interval (CI); ordered from oldest (top) to newest (bottom) study.

the study. Note that these findings describe the importance of different food items relative to other foods and cannot be used to infer the importance of food exposures compared to non-food exposures (e.g. environmental, animal, person-to-person exposures). Additionally, because we did not identify any case-control studies from the African, South-East Asian, nor Eastern Mediterranean subregions, the relevance of these results for these regions is unclear. Future investigation to determine the burden of foodborne STEC in these regions is needed.

As expected, this study corroborates the existing body of evidence that beef is a major source of exposure to STEC, thereby reinforcing the need for interventions in this commodity. Chicken was also identified as a significant food source, specifically in the Western Pacific A subregion. This finding differs from that of a recent global expert elicitation study [42], which ranked beef as the top food source of STEC infection in this subregion. However, it is important to note that the expert elicitation did not ask explicitly about poultry as a source of STEC (instead

Table 5. Results of the meta-analysis for each World Health Organization (WHO) Sub- Region, showing pooled univariate odds ratios (ORs) per food category (significant values shown in bold)

Food Category	WHO Subregion AMR A ^a (10 studies)			WHO Subregion AMR B ^a (one study)			WHO Subregion EUR A ^a (seven studies)			WHO Subregion WPR A ^a (three studies)		
	No. items per category (No. studies)	OR (95% C.I.)	I ²	No. items per category (No. studies)	OR (95% C.I.)	I ²	No. items per category (No. studies)	OR (95% C.I.)	I ²	No. items per category (No. studies)	OR (95% C.I.)	I ²
Beef	22 (9)	1.548 (1.086–2.207)^{b, c}	81%	32 (1)	1.555 (1.173–2.063)^{b, c}	69%	19 (6)	1.429 (1.044–1.956)	67%	7 (2)	1.243 (0.730–2.118)	62%
Meat – unspecified	9 (4)	1.545 (1.033–2.310)	64%	8 (1)	0.518 (0.380–0.704)^{b, c}	25%	38 (5)	1.172 (0.988–1.391) ^b	36%	5 (3)	1.295 (0.891–1.882)	0%
Produce	9 (3)	0.520 (0.369–0.734)^c	59%	0 (–)	N/A	N/A	17 (5)	0.872 (0.658–1.158)	52%	12 (3)	0.476 (0.188–1.206) ^b	89%
Dairy	1 (1)	9.774 (0.981–97.360)	0%	0 (–)	N/A	N/A	20 (7)	0.670 (0.507–0.886)	69%	2 (1)	1.209 (0.695–2.101)	0%
Chicken	4 (4)	0.335 (0.221–0.507)^c	0%	0 (–)	N/A	N/A	2 (2)	1.320 (0.170–10.273)	91%	3 (2)	2.689 (1.357–5.326)^c	0%
Seafood	2 (1)	0.683 (0.417–1.118)	0%	0 (–)	N/A	N/A	5 (2)	0.932 (0.385–2.258)	80%	1 (1)	0.452 (0.295–0.693)^c	0%
Pork	2 (2)	1.430 (0.841–2.431)	0%	2 (1)	1.107 (0.320–3.830)	76%	0 (–)	N/A	N/A	3 (2)	0.527 (0.379–0.733)^b	0%
Eggs	0 (–)	N/A	N/A	0 (–)	N/A	N/A	1 (1)	0.675 (0.477–0.956)^c	0%	4 (1)	0.642 (0.455–0.907)^c	0%
Lamb	0 (–)	N/A	N/A	0 (–)	N/A	N/A	3 (1)	1.936 (0.582–6.441)	45%	0 (–)	N/A	N/A
Turkey	1 (1)	0.347 (0.069–1.755)	0%	0 (–)	N/A	N/A	0 (–)	N/A	N/A	1 (1)	4.667 (0.374–58.248)	0%
Poultry/Game – unspecified	0 (–)	N/A	N/A	0 (–)	N/A	N/A	2 (2)	0.411 (0.228–0.740)^c	32%	0 (–)	N/A	N/A

^aWHO Subregions comprise the following countries:

Region of the Americas A (AMR A): Canada, Cuba, USA.

Region of the Americas B (AMR B): Antigua and Barbuda, Argentina, Bahamas, Barbados, Belize, Brazil, Chile, Colombia, Costa Rica, Dominica, Dominican Republic, El Salvador, Grenada, Guyana, Honduras, Jamaica, Mexico, Panama, Paraguay, Saint Kitts and Nevis, Saint Lucia, Saint Vincent and the Grenadines, Suriname, Trinidad and Tobago, Uruguay, Venezuela (Bolivarian Republic of).

European Region A (EUR A): Andorra, Austria, Belgium, Croatia, Cyprus, Czech Republic, Denmark, Finland, France, Germany, Greece, Iceland, Ireland, Israel, Italy, Luxembourg, Malta, Monaco, Netherlands, Norway, Portugal, San Marino, Slovenia, Spain, Sweden, Switzerland, United Kingdom of Great Britain and Northern Ireland.

Western Pacific Region A (WPR A): Australia, Brunei Darussalam, Japan, New Zealand, Singapore.

^bUsing trim and fill method.

^cThese items remained significant when clustering by the study was accounted for.

Table 6. Results from the meta-regression analysis of the univariate moderating effects of study characteristics, by food category, with significant values shown in bold (food categories with <20 results are not shown)

Study Characteristic (no. items; no. studies)	Characteristic level (no. items; no. studies)	OR	95% C.I.	P-value	I ²	Adjusted R ²
Beef (n = 80; 18)						
WHO subregion	AMR A (22; 9; reference)	–	–	–	72%	0%
	AMR B (32; 1)	1.121	0.732–1.717	0.600		
	EUR A (19; 6)	0.840	0.522–1.351	0.473		
	WPR A (7; 2)	0.714	0.368–1.385	0.319		
Publication year	(80; 18)	1.019	0.995–1.044	0.128	71%	4.7%
Study population age	All (37; 14; reference)	–	–	–	71%	1.5%
	Adults (5; 2)	1.327	0.689–2.552	0.397		
	Children (38; 4)	1.336	0.941–1.898	0.105		
Food item status	Not raw (1; 1; reference)	–	–	–	54%	55%
	Raw or undercooked (33; 9)	2.404	0.892–6.483	0.083		
	Unknown (46; 14)	1.036	0.387–2.776	0.944		
Robin's I	1 (14; 4; reference)	–	–	–	71%	3.0%
	2 (61; 11)	1.389	0.903–2.137	0.135		
	4 (4; 2)	1.009	0.448–2.274	0.982		
	6 (1; 1)	4.002	0.702–22.804	0.118		
Believable findings	Yes (79; 17; reference)	–	–	–	72%	1.8%
	No (1; 1)	3.122	0.564–17.270	0.192		
Meat-unspecified (n = 60; 13)						
WHO subregion	AMR A (9; 4; reference)	–	–	–	38%	60%
	AMR B (8; 1)	0.345	0.216–0.549	<0.001^a		
	EUR A (38; 5)	0.915	0.645–1.298	0.619		
	WPR A (5; 3)	0.866	0.488–1.537	0.624		
Publication year	(60; 13)	0.999	0.978–1.021	0.935	62%	0%
Study population age	All (24; 9; reference)	–	–	–	58%	11%
	Adults (10; 2)	1.453	0.940–2.246	0.093		
	Children (26; 5)	0.921	0.651–1.302	0.641		
Food item status	Not raw (10; 3; reference)	–	–	–	59%	5.5%
	Raw or undercooked (13; 5)	1.577	0.959–2.595	0.073		
	Unknown (37; 11)	1.266	0.825–1.941	0.280		
Robin's I	1 (24; 2; reference)	–	–	–	62%	0%
	2 (29; 9)	0.868	0.610–1.235	0.431		
	4 (7; 2)	1.126	0.671–1.891	0.654		
	6 (0; 0)	N/A	N/A	N/A		
Believable findings	Yes (59; 12; reference)	–	–	–	56%	22%
	No (1; 1)	7.523	2.073–27.302	0.002^a		
Produce (n = 38; 11)						
WHO subregion	AMR A (9; 3; reference)	–	–	–	65%	21%
	AMR B (0; 0)	N/A	N/A	N/A		
	EUR A (17; 5)	1.707	1.010–2.884	0.046		
	WPR A (12; 3)	1.149	0.650–2.029	0.633		

(Continued)

Table 6. (Continued.)

Study Characteristic (no. items; no. studies)	Characteristic level (no. items; no. studies)	OR	95% C.I.	P-value	I ²	Adjusted R ²
Publication year	(38; 11)	0.976	0.939–1.014	0.212	70%	7.6%
Study population age	All (26; 9; reference)	–	–	–	73%	0%
	Adults (4; 2)	0.888	0.412–1.916	0.762		
	Children (8; 3)	1.077	0.585–1.982	0.812		
Food item status	Not raw (0; 0)	N/A	N/A	N/A	72%	0%
	Raw or undercooked (16; 7; reference)	–	–	–		
	Unknown (22; 9)	1.035	0.649–1.650	0.885		
Robin's I	1 (12; 4; reference)	–	–	–	71%	2.2%
	2 (26; 7)	0.841	0.523–1.352	0.475		
	4 (0; 0)	N/A	N/A	N/A		
	6 (0; 0)	N/A	N/A	N/A		
Believable findings	Yes (38; 11; reference)	–	–	–	72%	N/A
	No (0; 0)	N/A	N/A	N/A		
Dairy (n = 23; 9)						
WHO subregion	AMR A (1; 1; reference)	–	–	–	66%	22%
	AMR B (0; 0)	N/A	N/A	N/A		
	EUR A (20; 7)	0.068	0.006–0.807	0.033		
	WPR A (2; 1)	0.127	0.009–1.695	0.118		
Publication year	(23; 9)	0.954	0.909–1.000	0.051	67%	10%
Study population age	All (9; 5; reference)	–	–	–	60%	37%
	Adults (3; 2)	0.887	0.374–2.104	0.786		
	Children (11; 4)	0.580	0.349–0.964	0.035		
Food item status	Not raw (2; 2; reference)	N/A	N/A	N/A	63%	10%
	Raw or undercooked (9; 6)	1.183	0.485–2.887	0.711		
	Unknown (12; 4)	0.557	0.247–1.254	0.157		
Robin's I	1 (7; 3; reference)	–	–	–	69%	1.2%
	2 (15; 5)	0.638	0.350–1.163	0.143		
	4 (1; 1)	10.040	0.789–127.810	0.076		
	6 (0; 0)	N/A	N/A	N/A		
Believable findings	Yes (22; 8; reference)	–	–	–	68%	12%
	No (1; 1)	13.783	1.145–165.894	0.039		

^aThese items remained significant when clustering by the study was accounted for.

asking explicitly about beef, small ruminants' and pigs' meat, dairy, vegetables and fruits and nuts and including an 'other' category within which additional foods could be identified). The discrepancy between our results and those from the expert elicitation may relate to the types of foods explicitly investigated by case-control vs. expert elicitation studies. Another reason may be that the contributors to the expert elicitation study were identifying food exposures for all STEC infections, both outbreak and sporadic and large outbreaks of STEC associated with chicken have rarely been reported, though STEC, including STEC O157, have been isolated from chickens and their meat [43, 44]. To guide policy and prevention decisions, methods are needed that allow different types of evidence about sources of foodborne

infections (e.g. from meta-analyses of case-control studies, outbreak analyses, expert elicitations, risk assessments) to be synthesised. Finally, chicken as a significant food source in the Western Pacific Region A could be an artifact of the analysis; however, the possibility that it is a significant source of exposure in this region cannot be discounted, as regional differences in significance could arise from differences in consumption patterns or preferred preparation methods of the commodity.

Here, produce, dairy, eggs, poultry/game – unspecified were also significant but had ORs less than one. For the purpose of source attribution, we do not draw conclusions on factors associated with a statistically significant reduced risk of disease, in part because of the impact of bias inherent in individual case-

control studies and thus to the final meta-analysis. While this is true for all exposures and all data that originate from interviews with patients and controls, it is particularly important when making inferences on the protective effect of specific exposures, which may eventually also be routes for infection [16,17]. Thus, any ORs less than one are reported herein but not interpreted further.

The way in which sporadic case-control studies are conducted and reported influences the type of data available for analyses such as this one. First, although case-control questionnaires of sporadic enteric illnesses like STEC often include an extensive list of potential risk factors, such as a list of food items, the specific food items included are often those for which there is an established or suspected association with illness, so as not to overburden study participants. Second, of the extensive list of food items for which data are collected, many may not have results reported in the final publication, or they may be reported in aggregate. For example, in one study, the questionnaire included 16 questions about different types of meats consumed, while the paper reported numerical results in aggregate as 'eating meat' [39].


Finally, publication culture prioritises reporting significant over non-significant results. Here, we noticed that significant findings were more often reported with extractable numbers, whereas non-significant findings were often reported as text without numeric values (e.g. 'eating ground beef was not associated with infection in this study' [28]), so we could not include these findings in our analyses. We attempted to overcome the limited availability of raw data by back-calculating raw values from ORs and SEs. Nevertheless, our results are likely skewed towards those food items for which there was established evidence or a strong hypothesis of risk at the time of the study, as well as results for which there were statistically significant findings. Thus, we recommend that primary research authors follow standard reporting guidelines for observational studies [45] and that they make the questionnaires and raw data available (e.g. supplementary materials, data repositories). Some of the more recent studies included here did provide questionnaires and summary data as supplementary materials [39e.g.]; we commend this comprehensive reporting while reiterating the need for studies to follow standard reporting guidelines and make publically accessible their questionnaires and raw data.

There were only 237 individual measures resulting from 21 unique studies included in the analysis and the majority of data were for beef and for the Americas A region, indicating even more sparse data for other food groups and regions. Indeed, our results from the Americas B region come from one study in a single country, conducted in children 0–15 years old, suggesting that the results for this subregion, in particular, be interpreted with caution. Additionally, there was temporal variation in study periods by region: in the Americas, studies were predominantly conducted in the mid-1980s and 1990s; in Europe, the early 1990s to early 2000s and in the Western Pacific region, the 2000s to 2012. Our findings are thus subject to limitations common to such meta-analyses based on sparse data and it is possible that the subregion effect seen here may actually be due to changes in risks over time. We explored publication bias using common methods, however, these tests are typically only reliable if there are 10 or more studies and their applicability to observational studies is unclear due to the presence of other biases (e.g. confounding) [46]. Therefore, significant publication bias tests identified in this review may be due to study confounding or other biases rather than possible publication bias. There are also some limitations to the meta-regression analyses used to explore the

potential effects of study characteristics on our estimates. For example, the exploration of significant covariates through meta-regression becomes increasingly less powerful as sample sizes (i.e. the number of studies) decrease and false positive findings are possible with multiple covariates assessed across numerous outcome groups [47]. Finally, we identified a potential effect of study bias on the results for meat – unspecified, dairy and chicken and thus they should be interpreted with caution. The results for beef, on the other hand, were not found to be affected by study bias and thus we have higher confidence in these findings.

Despite these limitations, our findings corroborate that interventions aimed at reducing the transmission of STEC via beef and other foods of animal origin, are important priorities in order to reduce the burden of foodborne STEC illness in the population, with additional studies needed to determine if this recommendation holds true for regions outside the Americas and Europe.

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References

- 1 Majowicz SE *et al.* (2014) Global incidence of human Shiga toxin-producing *Escherichia coli* infections and deaths: a systematic review and knowledge synthesis. *Foodborne Pathogens and Disease* **11**, 447–455.
- 2 Kirk MD *et al.* (2015) World Health Organization estimates of the global and regional disease burden of 22 foodborne bacterial, protozoal, and viral diseases, 2010: a data synthesis. *PLoS Medicine* **12**, e1001921.
- 3 Pires SM *et al.* (2009) Attributing the human disease burden of foodborne infections to specific sources. *Foodborne Pathogens and Disease* **6**, 417–424.
- 4 Kintz E *et al.* (2017) Transmission pathways for sporadic Shiga-toxin producing *E. coli* infections: a systematic review and meta-analysis. *International Journal of Hygiene and Environmental Health* **220**, 57–67.

- 5 Sargeant JM and O'Connor AM (2014) Introduction to systematic reviews in animal agriculture and veterinary medicine. *Zoonoses and Public Health* **61**, 3–9.
- 6 Hooman N *et al.* (2016) The prevalence of Shiga toxin-producing *Escherichia coli* in patients with gastroenteritis and sources of infections in Iran: a systematic review study protocol. *Journal of Pediatric Nephrology* **4**, 82–85.
- 7 Paudyal N *et al.* (2017) Prevalence of foodborne pathogens in food from selected African countries – a meta-analysis. *International Journal of Food Microbiology* **249**, 35–43.
- 8 Viswanathan M and Berkman ND (2012) Development of the RTI item bank on risk of bias and precision of observational studies. *Journal of Clinical Epidemiology* **65**, 163–178.
- 9 Viswanathan M *et al.* Assessing risk of bias and confounding in observational studies of interventions or exposures: further development of the RTI Item Bank. Methods Research Report. (Prepared by RTI-UNC Evidence-based Practice Center under Contract No. 290-2007-10056-I). AHRQ Publication No. 13-EHC106-EF. Rockville, MD: Agency for Healthcare Research and Quality; August 2013.
- 10 Sterne JAC (2016) ROBINS-I: a tool for assessing risk of bias in non-randomised studies of interventions. *British Medical Journal* **355**, i4919. doi: <https://doi.org/10.1136/bmj.i4919>.
- 11 Werber D *et al.* (2007) Shiga toxin-producing *Escherichia coli* infection in Germany - different risk factors for different age groups. *American Journal of Epidemiology* **165**, 425–434.
- 12 Richardson LC *et al.* (2017) An updated scheme for categorizing foods implicated in foodborne disease outbreaks: a tri-agency collaboration. *Foodborne Pathogens and Disease* **14**, 701–710.
- 13 Begg CB and Mazumdar M (1994) Operating characteristics of a rank correlation test for publication bias. *Biometrics* **50**, 1088–1101.
- 14 Egger M *et al.* (1997) Bias in meta-analysis detected by a simple, graphical test. *British Medical Journal* **315**, 629–634.
- 15 Duval S and Tweedie R (2000) Trim and fill: a simple funnel-plot-based method of testing and adjusting for publication bias in meta-analysis. *Biometrics* **56**, 455–463.
- 16 Domingues AR *et al.* (2012) Source attribution of human campylobacteriosis using a meta-analysis of case-control studies of sporadic infections. *Epidemiology and Infection* **140**, 970–981.
- 17 Domingues AR *et al.* (2012) Source attribution of human salmonellosis using a meta-analysis of case-control studies of sporadic infections. *Epidemiology and Infection* **140**, 959–969.
- 18 Viechtbauer W (2010) Conducting meta-analyses in R with the metafor package. *Journal of Statistical Software* **36**, 1–48.
- 19 MacDonald KL *et al.* (1988) *Escherichia coli* O157:H7, an emerging gastrointestinal pathogen. Results of a one-year, prospective, population-based study. *Journal of the American Medical Association* **259**, 3567–3570.
- 20 Bryant HE, Athar MA and Pai CH (1989) Risk factors for *Escherichia coli* O157:H7 infection in an urban community. *Journal of Infectious Diseases* **160**, 858–864.
- 21 Le Saux N *et al.* (1993) Ground beef consumption in noncommercial settings is a risk factor for sporadic *Escherichia coli* O157:H7 infection in Canada. *Journal of Infectious Diseases* **167**, 500–502.
- 22 Rowe PC *et al.* (1993) Diarrhoea in close contacts as a risk factor for childhood haemolytic uraemic syndrome. *Epidemiology and Infection* **110**, 9–16.
- 23 Slutsker L *et al.* (1998) A nationwide case-control study of *Escherichia coli* O157:H7 infection in the United States. *Journal of Infectious Diseases* **177**, 962–966.
- 24 Holton D *et al.* (1999) A Canadian multicentre case-control study of sporadic *Escherichia coli* O157:H7 infection. *Canadian Journal of Infectious Diseases* **10**, 117–121.
- 25 Finelli *et al.* (1995) Enhanced detection of sporadic *Escherichia coli* O157:H7 infections—New Jersey, July 1994. *Journal of the American Medical Association* **274**, 17–18.
- 26 Mead PS *et al.* (1997) Risk factors for sporadic infection with *Escherichia coli* O157:H7. *Archives of Internal Medicine* **157**, 204–208.
- 27 Parry SM *et al.* (1998) Risk factors for and prevention of sporadic infections with vero cytotoxin (shiga toxin) producing *Escherichia coli* O157. *Lancet* **351**, 1019–1022.
- 28 O'Brien SJ, Adak GK and Gilham C (2001) Contact with farming environment as a major risk factor for Shiga toxin (Vero cytotoxin)-producing *Escherichia coli* O157 infection in humans. *Emerging Infectious Diseases* **7**, 1049–1051.
- 29 Kassenborg HD *et al.* (2004) Farm visits and undercooked hamburgers as major risk factors for sporadic *Escherichia coli* O157:H7 infection: data from a case-control study in 5 FoodNet sites. *Clinical Infectious Diseases* **38**, S271–S278.
- 30 Piérard D *et al.* (1999) A case-control study of sporadic infection with O157 and non-O157 verocytotoxin-producing *Escherichia coli*. *Epidemiology and Infection* **122**, 359–365.
- 31 Locking ME *et al.* (2001) Risk factors for sporadic cases of *Escherichia coli* O157 infection: the importance of contact with animal excreta. *Epidemiology and Infection* **127**, 215–220.
- 32 Voetsch AC *et al.* (2007) Risk factors for sporadic Shiga toxin-producing *Escherichia coli* O157 infections in FoodNet sites, 1999–2000. *Epidemiology and Infection* **135**, 993–1000.
- 33 Vaillant V *et al.* (2009) Undercooked ground beef and person-to-person transmission as major risk factors for sporadic hemolytic uraemic syndrome related to Shiga-toxin producing *Escherichia coli* infections in children in France. *Pediatric Infectious Diseases Journal* **28**, 650–653.
- 34 Rivas M *et al.* (2008) Risk factors for sporadic Shiga toxin-producing *Escherichia coli* infections in children, Argentina. *Emerging Infectious Diseases* **14**, 763–771.
- 35 Hundy RL and Cameron S (2004) Risk factors for sporadic human infection with Shiga toxin-producing *Escherichia coli* in South Australia. *Communicable Disease Intelligence* **28**, 74–79.
- 36 Denno DM *et al.* (2009) Tri-county comprehensive assessment of risk factors for sporadic reportable bacterial enteric infection in children. *Journal of Infectious Diseases* **199**, 467–476.
- 37 McPherson M *et al.* (2009) Serogroup-specific risk factors for Shiga toxin-producing *Escherichia coli* infection in Australia. *Clinical Infectious Diseases* **49**, 249–256.
- 38 Friesema IH *et al.* (2015) Risk factors for sporadic Shiga toxin-producing *Escherichia coli* O157 and non-O157 illness in the Netherlands, 2008–2012, using periodically surveyed controls. *Epidemiology and Infection* **143**, 1360–1367.
- 39 Jaros P *et al.* (2013) A prospective case-control and molecular epidemiological study of human cases of Shiga toxin-producing *Escherichia coli* in New Zealand. *BMC Infectious Diseases* **13**, 1.
- 40 Kemp R (2005) The Epidemiology of VTEC O157, Non-O157 VTEC and Campylobacter spp. in a 100 km² Dairy Farming Area in Northwest England (PhD thesis). University of Liverpool, Liverpool, UK. Available at <http://www.opengrey.eu/item/display/10068/926085>.
- 41 Espie E (2007) Epidemiology of Shiga-Toxin Producing *Escherichia coli* Infections in Human in France (PhD thesis). Université Claude Bernard, Lyon, France. Available at <http://www.opengrey.eu/item/display/10068/814776>.
- 42 Hoffman S *et al.* (2017) Attribution of global foodborne disease to specific foods: findings from a World Health Organization structured expert elicitation. *PloS ONE* **12**, e0183641.
- 43 Persad A and Lejeune J (2015) Animal reservoirs of Shiga toxin-producing *Escherichia coli*. In Sperandio V and Hovde C (eds), *Enterohemorrhagic Escherichia coli and Other Shiga Toxin-Producing E. coli*. Washington, DC: ASM Press, pp. 231–244. doi: 10.1128/microbiolspec.EHEC-0027-2014.
- 44 Ferens WA and Hovde CJ (2011) *Escherichia coli* O157:H7: animal reservoir and sources of human infection. *Foodborne Pathogens and Disease* **8**, 465–487.
- 45 von Elm E *et al.* (2007) The Strengthening the Reporting of Observational Studies in Epidemiology (STROBE) statement: guidelines for reporting observational studies. *Bulletin of the World Health Organization* **85**, 867–872.
- 46 Sterne JAC *et al.* (2011) Recommendations for examining and interpreting funnel plot asymmetry in meta-analyses of randomised controlled trials. *The British Medical Journal* **343**, d4002.
- 47 Thompson SG and Higgins JP (2002) How should meta-regression analyses be undertaken and interpreted? *Statistics in Medicine* **21**, 1559–1573.